

In the Specification:

Please amend the specification as shown:

Please delete the paragraph on page 27, lines 22-28 and replace it with the following paragraph:

pMST73. The non-repeat region of FVO *glurp* was amplified with the primers 5'-CCC AGA TCT ACA AGT GAG AAT AGA AAT AAA C (SEQ ID NO: 8) (nucleotides 79 to 100, counting from A in the ATG start codon of the sequence of GenBank accession number M59706) and 5'-CCC AGA TCT TGC TTC ATG CTC GCT TTT TT CCG AT (SEQ ID NO: 9) (nucleotides 1475 to 1500 of the sequence of GenBank accession number M59706); digested with *Bgl*II, and the resulting DNA fragment was cloned into *Bgl*II digested pPSM1013.

Please delete the paragraph on page 28, lines 4-8 and replace it with the following paragraph:

pKBR7. The non-repeat region of F32 *glurp* was amplified with the primers 5'-AAG TAG ATC TAC TAA TAC AAG TGA GAA TAG AAA TAA AC (SEQ ID NO: 10) (nucleotides 73 to 100), and 5'-GTT CAG ATC TTT ATT CAT GAT GGC CTT CTA GC (SEQ ID NO: 11) (nucleotides 1519 to 1542); the resulting DNA fragment digested with *Bgl*II and cloned into *Bgl*II digested pPSM1013.

Please delete the paragraph on page 28, lines 11-18 and replace it with the following paragraph:

pKBR9. The C-terminal region of F32 *MSP3* was amplified with the primers 5'-CCC AGA TCT AAA GCA AAA GAA GCT TCT AGT TAT (SEQ ID NO: 12) (nucleotides 628 to 651) and 5'-ATT AGA TCT CAT TTA ATG ATT TTT AAA ATA TTT GGA TA (SEQ ID NO: 13), (nucleotides 1118 to 1140, counting from A in the ATG start codon of L07944); the resulting DNA fragment was digested with *Bgl*II and cloned into *Bgl*II digested pPSM1013. This *MSP3* region is identical to that of the FC27 allele (Genbank accession number L07944) except for the following residues at variable positions in *MSP3*: 735 (T→C) and 948 (A→G).